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FEB 21 2002  
TECH CENTER 1600/2900  
ENTERED

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/006,911

DATE: 12/17/2001

TIME: 10:45:25

Input Set : A:\RTS-0200 Sequence Listing.txt

Output Set: N:\CRF3\12172001\J006911.raw

6 <110> APPLICANT: William Gaarde  
7 Andrew T. Watt  
9 <120> TITLE OF INVENTION: ANTISENSE MODULATION OF COLLAPSin RESPONSE MEDIATOR PROTEIN  
2 EXPRESSION  
11 <130> FILE REFERENCE: RTS-0200  
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/006,911  
C--> 13 <141> CURRENT FILING DATE: 2001-11-08  
13 <160> NUMBER OF SEQ ID NOS: 89  
16 <210> SEQ ID NO: 1  
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18 <212> TYPE: DNA  
19 <213> ORGANISM: Artificial Sequence  
21 <220> FEATURE:  
23 <223> OTHER INFORMATION: Antisense Oligonucleotide  
25 <400> SEQUENCE: 1  
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31 <212> TYPE: DNA  
32 <213> ORGANISM: Artificial Sequence  
34 <220> FEATURE:  
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38 <400> SEQUENCE: 2  
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43 <211> LENGTH: 5421  
44 <212> TYPE: DNA  
45 <213> ORGANISM: Homo sapiens  
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49 tgactgaaaa aaatgactag ttattatgaa gacactactg ttgaagatgg atattttaac 120  
50 atggagtttc aacaaaatta cttcttgaga cagagctgat gtgtttttta aataacgtga 180  
51 ttttaagcat atatttgaac aaaactaaaa catttagtat tatgaatatg aaaaaagatc 240  
52 agtaaatcaa tgtactcttc taggctgaat taaggtagac tatttaaggt ttcaaaaaag 300  
53 tttggctggg gcagaataag ttttcaaaaa cccatgccat ccaaaattaa gatgacatgt 360  
54 agcagcaaga agtattccaa tgtctcataa ccagttctcg caagcaatgt gtattcctta 420  
55 ctttaaggaa gtgtcaaaaca aatagaaaaa tctggaagaa tttactaagt gtaataaatt 480  
56 agaggtaaat cgtataaaaa gaatttatgt ctacacaaaa tattcacaag tgggagtttt 540  
57 cttttacca cttctcagag tcttctagc cccctcttca cttctgaaag atgggattta 600  
58 ccaaaatctg gtttacattt aacttttcag ggacacatga cctgaaaaga aagatgtcag 660  
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60 gaattgggtc caaatggcat agaatacaac attatgtatc atgccaaata ccacttcctg 780  
61 cccaacaaaa ttcatcttt ctccagtaat gaagaggtgg acattcttgt tggactgtag 840  
62 catctgtgcc gccgcctcca caccaaccac ggcagctaac ctctgggcat catatttgga 900  
63 gtagagaaca gtgcagggtc acgtggcctc ttctcctctg ttgggtggctc tcagcatatt 960  
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66 ctctctctct ctctctctct tttttttccg ccctagctgg ggctgtgttg gaggagagga 1140

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67 agaaagagag acagaggatt gcattcatcc gttacgttct tgaaatttcc taatagcaag 1200
68 accagcgaag cggttgcacc cttttcaatc ttgcaaagga aaaaaacaaa acaaaacaaa 1260
69 aaaaacccaa gtcccttccc cggcagtttt tgccctaaag ctgccctctt gaaattaatt 1320
70 ttttcccagg agagagatgt cttatcaggg gaagaaaaat attccacgca tcacgagcga 1380
71 tegtcttctg atcaaaggag gtaaaattgt taatgatgac cagtcgttct atgcagacat 1440
72 atacatggaa gatgggttga tcaagcaaat aggagaaaaat ctgattgtgc caggaggagt 1500
73 gaagaccatc gaggccact cccggatggt gatccccgga ggaattgacg tccacactcg 1560
74 tttccagatg cctgatcagg gaatgacgtc tgetgatgat ttcttccaag gaaccaaggc 1620
75 ggccttggtt gggggaacca ctatgatcat tgaccacgtt gtctctgacg ctgggacaag 1680
76 cctgctcgct gcctttgacc agtggaggga atgggcccgc agcaagtcct gctgtgacta 1740
77 ctctctgcat gtggacatca gcgagtggca taagggcac caggaggaga tggaaagcgt 1800
78 tgtgaaggat cacggggtaa attccttccc cgtgtacatg gctttcaaag atcgcttcca 1860
79 gctaacggat tgccagattt atgaagtact gagtgtgatc cgggatattg gcgcatagc 1920
80 ccaagtcacac gcagaaaatg ggcacatcat tgcagaggag cagcagagga tccctggatc 1980
81 gggcatcacg ggccccgagg gacatgtgct gagccgacct gaggaggteg aggcgaagc 2040
82 cgtgaatcgt gccatcacca tgcacaacca gaaccaactgc ccgctgtata tcaccaaggt 2100
83 gatgagcaaa agctctgctg aggtcatcgc ccaggcacgg aagaaggga ctgtggtgta 2160
84 tggcagaccc atcactgcca gcttgggaac ggaaggctcc cattaactga gcaagaactg 2220
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89 ggatgagaac cagtttgttg ctgtgaccag caccaatgca gccaaagtct tcaaccttta 2520
90 cccccgaaa ggccgcattg ctgtgggac cgtatccgac ctggtcatct gggaccccga 2580
91 cagcgttaaa accatctctg ccaagacaca caacagctct ctcgagtaca acatctttga 2640
92 aggcattggg tgccgcgct cccactggt ggtcatcagc cagggggaaga ttgtcctgga 2700
93 ggacggcacc ctgcatgtca ccgaaggctc tggacgctac attccccgga agcccttccc 2760
94 tgattttgtt tacaagcgta tcaaggcaag gacagggctg gctgagctga gagggttcc 2820
95 tegtggcctg tatgacggac ctgtgtgtga agtgtctgtg acgcccaga cagtcactcc 2880
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102 ttttgetcat ccacttccc acacatctat gggatcaca cccaagacta cccaccaagc 3300
103 tcatacaggg aaccacaccc aacacttaga catgcgaaca agcagcccc agcgagggtc 3360
104 tcttctgct tcaacctcct agtgtctgtt agcattcctt ttcattgggg gagggagat 3420
105 aaagtgaatt gccagagct gcctttttct tttcttttta aaaattttta gaagttttcc 3480
106 ttgtggggtt ggggaggggc cggggtcagg gagagtcttt ttttttttt ttttaaatac 3540
107 taaattggaa catttaatte catattaata caagggttt gaactggaca tccaatgat 3600
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111 ctcttctctt ttaccttttt tctgctgtct ctcactctct ctttctctct ctagcttttt 3840
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113 tcggagacgg tgttttctct ccttgcctca ttatcttttc acctcccagg tctacatttc 3960
114 atggtggtcg ttgggtccgc cttaaaggatt tgagcgtttg ccattgcaag catagtgtcg 4020
115 tgtcatcctg gtccatgtag gactggtgct aaccacctgc catcatgagg atgtgtgcta 4080

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116 gagtgtggga ccttggecaa gtgcaggaat gggccatgcc gtctcaccga cagtatacaca 4140
117 cgtggaaccg cagacagggc ccagaagctt tagaggatg aggctgcaga accggagaga 4200
118 ttttccctctg tgcagtgtc tctggctaaa gtcacggtea aacctaaaca ccgagcctca 4260
119 ttaacccaag tgaaccaacc aaagtcacca gttcagaagt gctaagctaa taggagtctg 4320
120 acccgagggc ctgctgtctc ctggttaagt atcttttgag attctagaac acatgggagc 4380
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122 acatagaaag aggcctata aactcaaaaa gtcattggga aacttaaagt ctattctact 4500
123 ttgccaagag gagaaatgtg ttttatgaac gatagatcac atcagaactc ctgtggggag 4560
124 gaaaccttat aaattaaaca catggccccc ttagagacca caggcgatgt ctgtctccat 4620
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128 aagtaggtca ttccatcacc acccttgtct ctctacacat tttgccttg gggatctggt 4860
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131 aatttgaaca agttggaaaa aaacaatttt tgtttcaatt ctaagaaaca cttgcagctc 5040
132 tagtattcac ttgagctctc ctgtttttcc tgtaccgggt catggttaatt tttggttgtt 5100
133 ttggttgttt tcttaaaaaa caagttaaaa cctgacgatt tctgcagtga cttgatgtc 5160
134 taaaacagtg taggatttaa gaatagatgg tttttaatcc tggaaattgt gattgtgacc 5220
135 catgagtggg ggaactttca gttctaaagc tgataaagtg ttagccaga agagtacttt 5280
136 tttttgttaa ccactgtctt gatggcaaaa taattatggt aaaaaacaag tctcgtgttt 5340
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141 &lt;210&gt; SEQ ID NO: 4

142 &lt;211&gt; LENGTH: 21

143 &lt;212&gt; TYPE: DNA

144 &lt;213&gt; ORGANISM: Artificial Sequence

146 &lt;220&gt; FEATURE:

148 &lt;223&gt; OTHER INFORMATION: PCR Primer

150 &lt;400&gt; SEQUENCE: 4

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21

154 &lt;210&gt; SEQ ID NO: 5

155 &lt;211&gt; LENGTH: 21

156 &lt;212&gt; TYPE: DNA

157 &lt;213&gt; ORGANISM: Artificial Sequence

159 &lt;220&gt; FEATURE:

161 &lt;223&gt; OTHER INFORMATION: PCR Primer

163 &lt;400&gt; SEQUENCE: 5

164 qqgttaatga ggtcgggtg t

21

167 &lt;210&gt; SEQ ID NO: 6

168 &lt;211&gt; LENGTH: 30

169 &lt;212&gt; TYPE: DNA

170 &lt;213&gt; ORGANISM: Artificial Sequence

172 &lt;220&gt; FEATURE:

174 &lt;223&gt; OTHER INFORMATION: PCR Probe

176 &lt;400&gt; SEQUENCE: 6

177 cagtgtctctc tggctaaagt cacggcaca

30

180 &lt;210&gt; SEQ ID NO: 7

181 &lt;211&gt; LENGTH: 19

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Input Set : A:\RTS-0200 Sequence Listing.txt

Output Set: N:\CRF3\12172001\J006911.raw

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183 <213> ORGANISM: Artificial Sequence
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196 <213> ORGANISM: Artificial Sequence
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207 <211> LENGTH: 20
208 <212> TYPE: DNA
209 <213> ORGANISM: Artificial Sequence
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230 tgtgtgtgca tacatgtaca caattcccaa gttctgacca ctgagcctgg gatgagtggc 180
231 accacaatag taagcaactcc aatctaacct tttataataa tttcctacta aaagcaagca 240
232 gggctggagt agagaaacct ggaatgtctc attgtcaaaa aatacagcag tgcccaagta 300
233 gatgggggac atgtcaaaaag tacacagggg ctaacctgaa ggtgctccca gcggccatat 360
234 ttgagaccag ttacgcattc aaagaaataa tgatgtaaca aaccgtaacc cattgaataa 420
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241 aagatggata ttttaacatg gagtttcaac aaaattactt cttgagacag agctgatgtg 840
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244 ttaaggtttc aaaaaagttt ggctgggggc gaataagttt tacaaaacct atgcccacca 1020
245 aaattaagat gacatgtagc agcaagaagt attccaatgt ctcataacca gttctcgcaa 1080
246 gcaatgtgta ttcccttact taaggaagtg tcaaacaaat agaaaaatct ggaagaatgt 1140
247 actaagtgta ataaattaga ggtaaatcgt aataaaagaa tttatgtctc acaaaaaatat 1200
248 tcacaagtgg gagttttctt ctaccaactt ctgagagtc ttctagcccc ctcttcaact 1260

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Input Set : A:\RTS-0200 Sequence Listing.txt

Output Set: N:\CRF3\12172001\J006911.raw

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250 gaaaagaaag atgtcagata atactgacat tgcctcatgc actttctttg tatcagtcct 1380
251 tcttctgtaa gtaatcagaa ttgggtccaa atggcataga atcaaacatt atgtatcatg 1440
252 ccaaatacca ctctctgccc aacaaaattt catttttctc cagtaatgaa gaggtggaca 1500
253 ttcttggttg actgtagcat ctgtgcccgc cgctccacac caaccacggc agctaacctc 1560
254 tgggcatcat atttggagta gagaacagtg cagggtccacg tggcctcttc tcctctgttg 1620
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256 aaaatcaggt tcctgatgcc ttttattgta tacctgtcag aattggaaag aaagagctat 1740
257 gtcaacttga gaaacaagtt ctagaaaaac tacctaggag ttgtacagag gcalagtcca 1800
258 tgcttggtgc tcctgtctaa tggagaaaaa cagtggcaca agtaacagaa caaaagttaa 1860
259 tctaaactga ggtggcaaca tatttgaatt ctttttatga caacatattc gaattcctaa 1920
260 tatttggtct aagaaaatga taaaataaat acacatttct tgatcctaca ctgggtaggc 1980
261 tgaggtatca tgggaaagaa 2000
264 <210> SEQ ID NO: 11
265 <211> LENGTH: 4459
266 <212> TYPE: DNA
267 <213> ORGANISM: Homo sapiens
269 <400> SEQUENCE: 11
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272 aaatttctta atagcaagac cagcgaagcg gttgcacctt tttcaatctt gcaaaggaaa 180
273 aaaacaaaac aaaacaaaaa aaacccaagt ccccttcccg gcagtttttg ccttaaagct 240
274 gccctcttga aattaatttt tcccaggag agagatgtct tatcagggga agaaaaatat 300
275 tccacgcata acgagcgata gtcttctgat caaaggaggt aaaattgtta atgatgacca 360
276 gtcgttctat gcagacatat acatggaaga tgggttgatc aagcaaatag gagaaaatct 420
277 gattgtgcca ggaggagtga agaccatcga ggcccactcc cggatggtga tccccggagg 480
278 aattgacgtc cacactcggt tccagatgcc tgatcaggga atgacgtctg ctgatgattt 540
279 ctccaagga accaaggcgg ccttggtctg gggaaccact atgatcattg accacgttgt 600
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281 caagtcctgc tgtgactact ctctgcatgt ggacatcagc gagtggcata agggcatcca 720
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299 tgagctgaga ggggttctct gtggcctgta tgacggacct gtgtgtgaag tgtctgtgac 1800
300 gccaagaca gtcactccag cctctctggc caagacgtct cctgccaaagc agcaggcccc 1860

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## VERIFICATION SUMMARY

DATE: 12/17/2001

PATENT APPLICATION: US/10/006,911

TIME: 10:45:26

Input Set : A:\RTS-0200 Sequence Listing.txt

Output Set: N:\CRF3\12172001\J006911.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application No  
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:1363 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:3  
L:1366 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:89  
L:1366 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:3  
L:1370 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:89  
L:1370 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:4  
L:1371 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:89  
L:1371 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:4